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OM protein - protein search, using sw model

Run on: September 12, 2005, 20:15:12 ; Search time 567 Seconds
(without alignments)
272.693 Million cell updates/sec

Title: US-10-734-698A-39

Perfect score: 2083

Sequence: 1 MAETFLFTSSEVNEGHPDKL.....GREDDPFTWEVVKLWKA 392

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1777461 seqs, 39431504 residues

Total number of hits satisfying chosen parameters: 1777461

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database :

- Published Applications AA:*
- 1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
 - 2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
 - 3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
 - 4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
 - 5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
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 - 9: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
 - 10: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
 - 11: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
 - 12: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
 - 13: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
 - 14: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
 - 15: /cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
 - 16: /cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*
 - 17: /cgn2_6/ptodata/1/pubpaa/US10E_PUBCOMB.pep:*
 - 18: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
 - 19: /cgn2_6/ptodata/1/pubpaa/US11A_PUBCOMB.pep:*
 - 20: /cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
 - 21: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
 - 22: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2083	100.0	392	15	US-10-424-599-220046
2	2083	100.0	392	16	US-10-734-698A-39
3	2083	100.0	392	17	US-10-917-602A-39
4	2083	100.0	395	15	US-10-425-114-44212
5	2083	100.0	395	15	US-10-425-114-44833
6	2074	99.6	421	15	US-10-425-114-55424
7	2073	99.5	413	15	US-10-425-114-55057
8	2073	99.5	414	15	US-10-425-114-43817
9	2073	99.5	416	15	US-10-425-114-45713
10	2073	99.5	420	15	US-10-425-114-45878
11	2073	99.5	420	15	US-10-425-114-55959

12	2073	99.5	421	15	US-10-425-114-45874	Sequence 45874, A
13	2073	99.5	421	15	US-10-425-114-46284	Sequence 46284, A
14	2073	99.5	421	15	US-10-425-114-51724	Sequence 51724, A
15	2073	99.5	421	15	US-10-425-114-71917	Sequence 71917, A
16	2073	99.5	422	15	US-10-425-114-51415	Sequence 51415, A
17	2073	99.5	423	15	US-10-425-114-43754	Sequence 43754, A
18	2073	99.5	423	15	US-10-425-114-44212	Sequence 44212, A
19	2073	99.5	423	15	US-10-425-114-45712	Sequence 45712, A
20	2073	99.5	423	15	US-10-425-114-45723	Sequence 45723, A
21	2073	99.5	423	15	US-10-425-114-51430	Sequence 51430, A
22	2073	99.5	423	15	US-10-425-114-53367	Sequence 53367, A
23	2073	99.5	423	15	US-10-425-114-68219	Sequence 68219, A
24	2073	99.5	423	15	US-10-425-114-71903	Sequence 71903, A
25	2073	99.5	423	15	US-10-425-114-71916	Sequence 71916, A
26	2073	99.5	423	15	US-10-425-114-71922	Sequence 71922, A
27	2073	99.5	423	15	US-10-425-114-71923	Sequence 71923, A
28	2073	99.5	423	15	US-10-425-114-71929	Sequence 71929, A
29	2073	99.5	423	15	US-10-425-114-71954	Sequence 71954, A
30	2073	99.5	423	15	US-10-425-114-71966	Sequence 71966, A
31	2073	99.5	423	15	US-10-425-114-71968	Sequence 71968, A
32	2073	99.5	423	15	US-10-425-114-71969	Sequence 71969, A
33	2073	99.5	423	15	US-10-425-114-71975	Sequence 71975, A
34	2073	99.5	423	15	US-10-425-114-71992	Sequence 71992, A
35	2073	99.5	423	15	US-10-425-114-71993	Sequence 71993, A
36	2073	99.5	423	15	US-10-425-114-71994	Sequence 71994, A
37	2073	99.5	423	15	US-10-425-114-72190	Sequence 72190, A
38	2073	99.5	425	15	US-10-425-114-45846	Sequence 45846, A
39	2062.5	99.0	393	15	US-10-424-599-220047	Sequence 220047, A
40	2056.5	98.7	393	15	US-10-424-599-258237	Sequence 258237, A
41	2048.5	98.3	393	15	US-10-424-599-220043	Sequence 220043, A
42	1999	96.0	395	15	US-10-424-599-260160	Sequence 260160, A
43	1999	96.0	395	15	US-10-424-599-260162	Sequence 260162, A
44	1999	96.0	402	15	US-10-425-114-44433	Sequence 44433, A
45	1999	96.0	420	15	US-10-425-114-44746	Sequence 44746, A

ALIGNMENTS

RESULT 1
US-10-424-599-220046
; Sequence 220046, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovacic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 220046
; LENGTH: 392
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_40730C.1.pep
US-10-424-599-220046

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Matches	392	Conservative	0	Mismatches	0	Indels	0
		Gaps	0				
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Db	1	MAETFLFTSSEVNEGHPDKL	CDQISDAVLADCL	EQDDPSKVCAC	TCTKTNLVVFG	GEIT	60
QY	61	KANVDYKIVTRTCR	NIQFVSNVDGLD	ADNCKVLVNI	EQQSPDIAQGV	HGHLTKR	PEETG 120
Db	61	KANVDYKIVTRTCR	NIQFVSNVDGLD	ADNCKVLVNI	EQQSPDIAQGV	HGHLTKR	PEETG 120

QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTEVRKNGTCPLWRPDGKTQVTVEYND 180
DB 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTEVRKNGTCPLWRPDGKTQVTVEYND 180
QY 181 NGAMVPRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
DB 181 NGAMVPRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
QY 241 IGGPHDAGLTGRKIIIDTYGGWGAHGGAGFGSKDPTKVDKRGAYIVRQAASIVASGLA 300
DB 241 IGGPHDAGLTGRKIIIDTYGGWGAHGGAGFGSKDPTKVDKRGAYIVRQAASIVASGLA 300
QY 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
DB 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
QY 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
DB 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392

RESULT 2
US-10-734-698A-39
; Sequence 39, Application US/10734698A
; Publication No. US20040209341A1
; GENERAL INFORMATION:
; APPLICANT: FALCO, SAVERIO CARL
; ALLEN, STEPHEN M.
; RAFALSKI, J. ANTONI
; HITZ, WILLIAM D.
; KINNEY, ANTHONY J.
; ABELL, LYNNE N.
; THORPE, CATHERINE J.
; TITLE OF INVENTION: PLANT AMINO ACID BIOSYNTHETIC ENZYMES
; NUMBER OF SEQUENCES: 43
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: E. I. DU PONT DE NEMOURS AND COMPANY
; STREET: 1007 MARKET STREET
; CITY: WILMINGTON
; STATE: DELAWARE
; COUNTRY: USA
; ZIP: 19898
; COMPUTER READABLE FORM:
; MEDIUM TYPE: DISKETTE, 3.50 INCH
; COMPUTER: IBM PC COMPATIBLE
; OPERATING SYSTEM: MICROSOFT WINDOWS 95
; SOFTWARE: MICROSOFT WORD VERSION 7.0A
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/734,698A
; FILING DATE: 12-Dec-2003
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/048,771
; FILING DATE: 6-Jun-1997
; ATTORNEY/AGENT INFORMATION:
; NAME: MAJARIAN, WILLIAM R.
; REGISTRATION NUMBER: 41,173
; REFERENCE/DOCKET NUMBER: BB-1087
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 302-992-4926
; TELEFAX: 302-773-0164
; INFORMATION FOR SEQ ID NO: 39:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 392 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; IMMEDIATE SOURCE:
; CLONE: 82.12B06
; SEQUENCE DESCRIPTION: SEQ ID NO: 39:
US-10-734-698A-39

Query Match 100.0%; Score 2083; DB 16; Length 392;
Best Local Similarity 100.0%; Pred. No. 9.1e-191;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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DB 1 MAETFLFTSESVNEGHDPDKLQDISDAVLDAQLDQDPDSKVACETCTKTNLVMVFEITTT 60
QY 61 KANVDYEKIVRDTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGLTKRPEEIG 120
DB 61 KANVDYEKIVRDTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGLTKRPEEIG 120
QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTEVRKNGTCPLWRPDGKTQVTVEYND 180
DB 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTEVRKNGTCPLWRPDGKTQVTVEYND 180
QY 181 NGAMVPRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
DB 181 NGAMVPRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
QY 241 IGGPHDAGLTGRKIIIDTYGGWGAHGGAGFGSKDPTKVDKRGAYIVRQAASIVASGLA 300
DB 241 IGGPHDAGLTGRKIIIDTYGGWGAHGGAGFGSKDPTKVDKRGAYIVRQAASIVASGLA 300
QY 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
DB 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
QY 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
DB 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392

RESULT 3
US-10-917-602A-39
; Sequence 39, Application US/10917602A
; Publication No. US20050120405A1
; GENERAL INFORMATION:
; APPLICANT: Falco, Saverio Carl
; APPLICANT: Liu, Zhan-Bin
; TITLE OF INVENTION: Plant Amino Acid Biosynthetic Enzymes
; FILE REFERENCE: BB-1087 US CJP
; CURRENT APPLICATION NUMBER: US/10/917,602A
; CURRENT FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: US 10/734698
; PRIOR FILING DATE: 2003-12-12
; PRIOR APPLICATION NUMBER: US 09/424978
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US98/11692
; PRIOR FILING DATE: 1998-06-05
; PRIOR APPLICATION NUMBER: 60/049,443
; PRIOR FILING DATE: 1997-06-12
; PRIOR APPLICATION NUMBER: US 60/048,771
; PRIOR FILING DATE: 1997-06-06
; NUMBER OF SEQ ID NOS: 69
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 39
; LENGTH: 392
; TYPE: PRT
; ORGANISM: Glycine max
US-10-917-602A-39

Query Match 100.0%; Score 2083; DB 17; Length 392;
Best Local Similarity 100.0%; Pred. No. 9.1e-191;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAETFLFTSESVNEGHDPDKLQDISDAVLDAQLDQDPDSKVACETCTKTNLVMVFEITTT 60
DB 1 MAETFLFTSESVNEGHDPDKLQDISDAVLDAQLDQDPDSKVACETCTKTNLVMVFEITTT 60
QY 61 KANVDYEKIVRDTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGLTKRPEEIG 120

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Db      61 KANVDYEKIVRDTCRNIGFVNSDVGDLADNCKVLVNIQQSPDIAQGVGHGHLTKRPEEIG 120
Qy      121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 180
Db      121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 180
Qy      181 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
Db      181 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
Qy      241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 300
Db      241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 300
Qy      301 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 360
Db      301 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 360
Qy      361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db      361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392

RESULT 4
US-10-425-114-44212
; Sequence 44212, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 44212
; LENGTH: 395
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 700896469_FLI.pep
US-10-425-114-44212

Query Match      100.0%; Score 2083; DB 15; Length 395;
Best Local Similarity 100.0%; Pred. No. 9.2e-191;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 MAETFLFTSESNEGHPDKLDCQISDAVLDAQLEQDPDSKVACETCTKTNLVMVFEIIT 60
Db      4 MAETFLFTSESNEGHPDKLDCQISDAVLDAQLEQDPDSKVACETCTKTNLVMVFEIIT 63
Qy      61 KANVDYEKIVRDTCRNIGFVNSDVGDLADNCKVLVNIQQSPDIAQGVGHGHLTKRPEEIG 120
Db      64 KANVDYEKIVRDTCRNIGFVNSDVGDLADNCKVLVNIQQSPDIAQGVGHGHLTKRPEEIG 123
Qy      121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 180
Db      124 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 183
Qy      181 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
Db      184 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 243
Qy      241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 300
Db      244 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 303
Qy      301 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 360
Db      301 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 360
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Db      304 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 363
Qy      361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db      364 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 395

RESULT 5
US-10-425-114-44833
; Sequence 44833, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 44833
; LENGTH: 395
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701150545_FLI.pep
US-10-425-114-44833

Query Match      100.0%; Score 2083; DB 15; Length 395;
Best Local Similarity 100.0%; Pred. No. 9.2e-191;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 MAETFLFTSESNEGHPDKLDCQISDAVLDAQLEQDPDSKVACETCTKTNLVMVFEIIT 60
Db      4 MAETFLFTSESNEGHPDKLDCQISDAVLDAQLEQDPDSKVACETCTKTNLVMVFEIIT 63
Qy      61 KANVDYEKIVRDTCRNIGFVNSDVGDLADNCKVLVNIQQSPDIAQGVGHGHLTKRPEEIG 120
Db      64 KANVDYEKIVRDTCRNIGFVNSDVGDLADNCKVLVNIQQSPDIAQGVGHGHLTKRPEEIG 123
Qy      121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 180
Db      124 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYND 183
Qy      181 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 240
Db      184 NGAMVPVRVHTVLISQHDSTVTNDEIAADLKEHVIKPVIPKYLDEKTIIFHLNPSGRFV 243
Qy      241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 300
Db      244 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKVDKSGAYIVRQAASIVASGLA 303
Qy      301 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 360
Db      304 RRCIVQVSAIGAIVPEPLSVFVDYTGTKIHDKIELNIVKENFDFRPGMISINLDLKRGN 363
Qy      361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db      364 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 395

RESULT 6
US-10-425-114-55424
; Sequence 55424, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
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; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 55424
; LENGTH: 421
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-GMFLMINSOY064D01_FLI.pep
US-10-425-114-55424

Query Match          99.6%; Score 2074; DB 15; Length 421;
Best Local Similarity 99.5%; Pred. No. 7.4e-190;
Matches 390; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 60
Db 30 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 89

QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 120
Db 90 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 149

QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVKRNKGTCPWLRPDGKTQVTVEYND 180
Db 150 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVKRNKGTCPWLRPDGKTQVTVEYND 209

QY 181 NGAMVPRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 240
Db 210 NGARVPVRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 269

QY 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGDKPTKVDKRGAYIVRQAASIVASGLA 300
Db 270 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGDKPTKVDKRGAYIVRQAASIVASGLA 329

QY 301 RRCIVQVSAIGAIVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
Db 330 RRCIVQVSAIGAIVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 389

QY 361 NRFLKTAAYGHFGREDPDPFTWEVVKPLKWEKA 392
Db 390 NRFLKTAAYGHFGREDPDPFTWEVVKPLKWEKA 421

RESULT 7
US-10-425-114-55057
; Sequence 55057, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 55057
; LENGTH: 413
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-GMFLMINSOY064D01_FLI.pep
US-10-425-114-55057

Query Match          99.5%; Score 2073; DB 15; Length 414;
Best Local Similarity 99.2%; Pred. No. 9e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 60
Db 23 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 82

QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 120
Db 83 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 142
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; OTHER INFORMATION: Clone ID: 701211701_FLI.pep
US-10-425-114-55057

Query Match          99.5%; Score 2073; DB 15; Length 413;
Best Local Similarity 99.2%; Pred. No. 8.9e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 60
Db 22 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 81

QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 120
Db 82 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 141

QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVKRNKGTCPWLRPDGKTQVTVEYND 180
Db 142 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVKRNKGTCPWLRPDGKTQVTVEYND 201

QY 181 NGAMVPRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 240
Db 202 NGARVPVRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 261

QY 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGDKPTKVDKRGAYIVRQAASIVASGLA 300
Db 262 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGDKPTKVDKRGAYIVRQAASIVASGLA 321

QY 301 RRCIVQVSAIGAIVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 360
Db 322 RRCIVQVSAIGAIVPEPLSVFVDTYGTGKIHDKEILNIVKENFDFRPGMISINLDLKRGN 381

QY 361 NRFLKTAAYGHFGREDPDPFTWEVVKPLKWEKA 392
Db 382 NRFLKTAAYGHFGREDPDPFTWEVVKPLKWEKA 413

RESULT 8
US-10-425-114-43817
; Sequence 43817, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 43817
; LENGTH: 414
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 700649684_FLI.pep
US-10-425-114-43817

Query Match          99.5%; Score 2073; DB 15; Length 414;
Best Local Similarity 99.2%; Pred. No. 9e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 60
Db 23 MAETFLFTSESNEGHDPDKLQDISDAVLDAACLEQDPDPSKVACETCTKTNLVMVFGEIT 82

QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 120
Db 83 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLVNIQQSPDIAQQVGHGHLTKRPEEIG 142
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QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 180
Db 143 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 202
QY 181 NGAMVPRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 240
Db 203 NGARVPIRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 262
QY 241 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 300
Db 263 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 322
QY 301 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 360
Db 323 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 382
QY 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db 383 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 414

RESULT 9
US-10-425-114-45713
; Sequence 45713, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 45713
; LENGTH: 416
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 700894196_FLI.pep
US-10-425-114-45713

Query Match 99.5%; Score 2073; DB 15; Length 416;
Best Local Similarity 99.2%; Pred. No. 9e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPKLCQISDAVLDALEQDPSKVACETCTKTNLVMVFGEIT 60
Db 25 MAETFLFTSESNEGHDPKLCQISDAVLDALEQDPSKVACETCTKTNLVMVFGEIT 84
QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSQSDIAQGVHGLTKRPEEIG 120
Db 85 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSQSDIAQGVHGLTKRPEEIG 144
QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 180
Db 145 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 204
QY 181 NGAMVPRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 240
Db 205 NGARVPIRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 264
QY 241 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 300
Db 265 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 324
QY 301 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 360
Db 325 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 384
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QY 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db 385 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 416

RESULT 10
US-10-425-114-45878
; Sequence 45878, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 45878
; LENGTH: 420
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701040251_FLI.pep
US-10-425-114-45878

Query Match 99.5%; Score 2073; DB 15; Length 420;
Best Local Similarity 99.2%; Pred. No. 9.2e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAETFLFTSESNEGHDPKLCQISDAVLDALEQDPSKVACETCTKTNLVMVFGEIT 60
Db 29 MAETFLFTSESNEGHDPKLCQISDAVLDALEQDPSKVACETCTKTNLVMVFGEIT 88
QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSQSDIAQGVHGLTKRPEEIG 120
Db 89 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSQSDIAQGVHGLTKRPEEIG 148
QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 180
Db 149 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGGKTQVTVYYND 208
QY 181 NGAMVPRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 240
Db 209 NGARVPIRVHTVLISQHDSTVNDDEIAADLKEHVIKPVPEKYLDEKTI FHLNPSGRFV 268
QY 241 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 300
Db 269 IGGPHGDAGLTGRKIIIDITYGGWGAHGGGAFSGKDPKVDKRSYAIIVROAAKSIIVASGLA 328
QY 301 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 360
Db 329 RRCIVQVSAIGAIVPEPLSVFVDITYGKIHDKIELNIVKENFDFRPGMISINLCLKRGN 388
QY 361 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 392
Db 389 NRFLKTAAYGHFGREDPDFTWEVVKPLKWEKA 420

RESULT 11
US-10-425-114-55959
; Sequence 55959, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
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RESULT 12
US-10-425-114-45874
; Sequence 45874, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kowalik, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(5313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 45874
; LENGTH: 421
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701001311_FLI.pep
US-10-425-114-45874

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Db 210 NGARVPIRVHTVLISQHDVTNDEIAADLKEHVIKPVIPEKYLDEKTI FHLNPSGRFV 269
Qy 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKPTKVDKRSYIVRQAASIVASGLA 300
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Qy 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHKDEILNI VKENFDFRPGMISINLDKRCGN 360
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Db 390 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 421

RESULT 14
US-10-425-114-51724
; Sequence 51724, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 51724
; LENGTH: 421
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 700961178_FLI.pap
US-10-425-114-51724

Query Match 99.5%; Score 2073; DB 15; Length 421;
Best Local Similarity 99.2%; Pred. No. 9.2e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MAETFLFTSESNEGHDPDKLDCQISDAVLDALEQDPPDSKVACETCTKTNLVMVFGEIIT 60
Db 30 MAETFLFTSESNEGHDPDKLDCQISDAVLDALEQDPPDSKVACETCTKTNLVMVFGEIIT 89
Qy 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVHGHILTKRPEEIG 120
Db 90 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVHGHILTKRPEEIG 149
Qy 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVBYND 180
Db 150 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVBYND 209
Qy 181 NGAMVPVRVHTVLISQHDVTNDEIAADLKEHVIKPVIPEKYLDEKTI FHLNPSGRFV 240
Db 210 NGARVPIRVHTVLISQHDVTNDEIAADLKEHVIKPVIPEKYLDEKTI FHLNPSGRFV 269
Qy 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKPTKVDKRSYIVRQAASIVASGLA 300
Db 270 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKPTKVDKRSYIVRQAASIVASGLA 329
Qy 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHKDEILNI VKENFDFRPGMISINLDKRCGN 360
Db 330 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHKDEILNI VKENFDFRPGMISINLDKRCGN 389
Qy 361 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db 390 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 421
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Job time : 569 secs

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Db 390 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 421

RESULT 15
US-10-425-114-71917
; Sequence 71917, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 71917
; LENGTH: 421
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 700751645_FLI.pap
US-10-425-114-71917

Query Match 99.5%; Score 2073; DB 15; Length 421;
Best Local Similarity 99.2%; Pred. No. 9.2e-190;
Matches 389; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MAETFLFTSESNEGHDPDKLDCQISDAVLDALEQDPPDSKVACETCTKTNLVMVFGEIIT 60
Db 30 MAETFLFTSESNEGHDPDKLDCQISDAVLDALEQDPPDSKVACETCTKTNLVMVFGEIIT 89
Qy 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVHGHILTKRPEEIG 120
Db 90 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVHGHILTKRPEEIG 149
Qy 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVBYND 180
Db 150 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVBYND 209
Qy 181 NGAMVPVRVHTVLISQHDVTNDEIAADLKEHVIKPVIPEKYLDEKTI FHLNPSGRFV 240
Db 210 NGARVPIRVHTVLISQHDVTNDEIAADLKEHVIKPVIPEKYLDEKTI FHLNPSGRFV 269
Qy 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKPTKVDKRSYIVRQAASIVASGLA 300
Db 270 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKPTKVDKRSYIVRQAASIVASGLA 329
Qy 301 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHKDEILNI VKENFDFRPGMISINLDKRCGN 360
Db 330 RRCIVQVSYAIGVPEPLSVFVDTYGTGKIHKDEILNI VKENFDFRPGMISINLDKRCGN 389
Qy 361 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db 390 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 421
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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: September 12, 2005, 14:28:42 ; Search time 30 Seconds
(without alignments)
975.414 Million cell updates/sec

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Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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6: /cgn2_6/ptodata/1/iaa/backfile1.pcp.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2083	100.0	392	4	US-09-424-978B-39 Sequence 39, Appl
2	1946	93.4	396	4	US-09-424-978B-36 Sequence 36, Appl
3	1891.5	90.8	394	4	US-09-424-978B-42 Sequence 42, Appl
4	1316.5	63.2	395	4	US-09-424-978B-471 Sequence 471, Appl
5	1316.5	63.2	416	4	US-09-949-016-10059 Sequence 10059, A
6	1280.5	61.5	390	4	US-09-248-796A-18255 Sequence 18255, A
7	1274.5	61.2	395	4	US-09-949-016-5939 Sequence 5939, Ap
8	1274.5	61.2	401	4	US-09-949-016-7658 Sequence 7658, Ap
9	1102	52.9	404	4	US-09-107-532A-6821 Sequence 6821, Ap
10	1086	52.1	387	4	US-09-543-681A-7130 Sequence 7130, Ap
11	1068.5	51.3	385	4	US-09-489-039A-11917 Sequence 11917, A
12	1062.5	51.0	396	4	US-09-583-110-2778 Sequence 2778, Ap
13	1062.5	51.0	405	4	US-09-107-433-2830 Sequence 2830, Ap
14	1061.5	51.0	396	3	US-09-273-686-2 Sequence 2, Appl
15	1055	50.6	395	4	US-09-328-352-6660 Sequence 6660, Ap
16	1042.5	50.0	415	3	US-09-134-001C-5077 Sequence 5077, Ap
17	1040	49.9	388	4	US-09-340-236-3444 Sequence 3444, Ap
18	1028.5	49.4	402	2	US-08-403-852D-19 Sequence 19, Appl
19	1028.5	49.4	402	3	US-08-510-646B-20 Sequence 20, Appl
20	1028.5	49.4	402	3	US-09-231-818-19 Sequence 19, Appl
21	1028.5	49.4	402	4	US-09-635-359B-19 Sequence 19, Appl
22	1006	48.3	407	3	US-08-955-957A-2 Sequence 2, Appl
23	1000.5	48.0	401	4	US-09-252-991A-19899 Sequence 19899, A
24	832	39.9	332	3	US-09-320-878-16 Sequence 16, Appl
25	832	39.9	332	4	US-09-141-908-20 Sequence 20, Appl
26	832	39.9	332	4	US-09-657-440-16 Sequence 16, Appl
27	803.5	38.6	313	4	US-09-902-540-10716 Sequence 10716, A

ALIGNMENTS

RESULT 1

US-09-424-978B-39

; Sequence 39, Application US/09424978B

; Patent No. 6664445

; GENERAL INFORMATION:

; APPLICANT: Falco, Saverio Carl

; APPLICANT: Allen, Stephen M.

; APPLICANT: Rafalski, J. Antoni

; APPLICANT: Hitz, William D.

; APPLICANT: Kinney, Anthony J.

; APPLICANT: Abell, Lynne N.

; APPLICANT: Thorpe, Catherine J.

; TITLE OF INVENTION: Plant Amino Acid Biosynthetic Enzymes

; FILE REFERENCE: BB-1087

; CURRENT APPLICATION NUMBER: US/09/424,978B

; CURRENT FILING DATE: 1999-12-02

; PRIOR APPLICATION NUMBER: US 60/048,771

; PRIOR FILING DATE: 1997-06-06

; NUMBER OF SEQ ID NOS: 43

; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 39

; LENGTH: 392

; TYPE: PRT

; ORGANISM: Glycine max

US-09-424-978B-39

Query Match 100.0%; Score 2083; DB 4; Length 392;

Best Local Similarity 100.0%; Pred. No. 4.1e-212; Mismatches 0; Indels 0; Gaps 0;

Matches 392; Conservative 0;

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Db 1 MAETFLFTSESVEGHPDKLQDSDAVLDALEQDDPSKVACETCTKTNLVWVFGEIT 60

QY 61 KANVDYEKIVRDTCTNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVGHULTKRPPEIG 120

Db 61 KANVDYEKIVRDTCTNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVGHULTKRPPEIG 120

QY 121 AGDQGHMGYATDETPELMPLSHVLATKLGARLTVRNKGTCPWLRRPGKQTQVVEYND 180

Db 121 AGDQGHMGYATDETPELMPLSHVLATKLGARLTVRNKGTCPWLRRPGKQTQVVEYND 180

QY 181 NGAMPVVRVHTVLSTQHDVTWNDEIAADLKEHVIKPVIPEKYLDKXTIHLNPSGRFV 240

Db 181 NGAMPVVRVHTVLSTQHDVTWNDEIAADLKEHVIKPVIPEKYLDKXTIHLNPSGRFV 240

QY 241 IGGPHGDAGLTKRKIIIDTYGSGWAGHGGAFSGKDPKTKVDRSGAYIVRQAASIVASGLA 300

Db 241 IGGPHGDAGLTKRKIIIDTYGSGWAGHGGAFSGKDPKTKVDRSGAYIVRQAASIVASGLA 300

QY 301 RRCIVQSVYAIGVPEPLSVFVDTYGTGKHDKHEILNIVKENFDFRPGMISINLDLKRGN 360
Db |||||
QY 301 RRCIVQSVYAIGVPEPLSVFVDTYGTGKHDKHEILNIVKENFDFRPGMISINLDLKRGN 360
Db |||||
QY 361 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db |||||
QY 361 NRFLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db |||||

RESULT 2
US-09-424-978B-36
; Sequence 36, Application US/09424978B
; Patent No. 6664445
; GENERAL INFORMATION:
; APPLICANT: Falco, Saverio Carl
; APPLICANT: Allen, Stephen M.
; APPLICANT: Rafaleki, J. Antoni
; APPLICANT: Hitz, William D.
; APPLICANT: Kinney, Anthony J.
; APPLICANT: Abell, Lynne N.
; APPLICANT: Thorpe, Catherine J.
; TITLE OF INVENTION: Plant Amino Acid Biosynthetic Enzymes
; FILE REFERENCE: BB-1087
; CURRENT APPLICATION NUMBER: US/09/424,978B
; CURRENT FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: US 60/048,771
; PRIOR FILING DATE: 1997-06-06
; NUMBER OF SEQ ID NOS: 43
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 36
; LENGTH: 396
; TYPE: PRT
; ORGANISM: Zea mays
US-09-424-978B-36

Query Match 93.4%; Score 1946; DB 4; Length 396;
Best Local Similarity 93.3%; Pred. No. 1.4e-197;
Matches 363; Conservative 14; Mismatches 12; Indels 0; Gaps 0;
QY 3 2TFLFTSESNEGHPDKLDCQISDAVLDAACLEQDPDSKVACETCTKTNLVMVFGIITKA 62
Db |||||
QY 5 DFLFTSESNEGHPDKLDCQISDAVLDAACLEQDPDSKVACETCTKTNLVMVFGIITKA 64
Db |||||
QY 63 NVDEYKIVRTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGHHTKRPBEIGAG 122
Db |||||
QY 65 NVDEYKIVRTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGHHTKRPBEIGAG 124
Db |||||
QY 123 DQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGKTQVTVVEYNDNG 182
Db |||||
QY 125 DQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGKTQVTVVEYNDNG 184
Db |||||
QY 183 AMVPVRVHTVLTSTQHDFTVNDIEAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFVIG 242
Db |||||
QY 185 AMVPVRVHTVLTSTQHDFTVNDIEAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFVIG 244
Db |||||
QY 243 GPHGDAGLTKRKIIIDTYGCGAHGGGAFSGKDPKVDKRGAYIVRQAAKSIIVASGLARR 302
Db |||||
QY 245 GPHGDAGLTKRKIIIDTYGCGAHGGGAFSGKDPKVDKRGAYIVRQAAKSIIVASGLARR 304
Db |||||
QY 303 CIQVQSVYAIGVPEPLSVFVDTYGTGKHDKHEILNIVKENFDFRPGMISINLDLKRGNR 362
Db |||||
QY 305 AIQVQSVYAIGVPEPLSVFVDTYGTGAIPDKELKIVKENFDFRPGMIIINLDLKRGNR 364
Db |||||
QY 363 FLKTAAYGHGREDPDFTWEVVKPLKWEK 391
Db |||||
QY 365 YLKTAAYGHGREDPDFTWEVVKPLKWEK 393
Db |||||

RESULT 3
US-09-424-978B-42
; Sequence 42, Application US/09424978B
; Patent No. 6664445
; GENERAL INFORMATION:

; APPLICANT: Falco, Saverio Carl
; APPLICANT: Allen, Stephen M.
; APPLICANT: Rafaleki, J. Antoni
; APPLICANT: Hitz, William D.
; APPLICANT: Kinney, Anthony J.
; APPLICANT: Abell, Lynne N.
; APPLICANT: Thorpe, Catherine J.
; TITLE OF INVENTION: Plant Amino Acid Biosynthetic Enzymes
; FILE REFERENCE: BB-1087
; CURRENT APPLICATION NUMBER: US/09/424,978B
; CURRENT FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: US 60/048,771
; PRIOR FILING DATE: 1997-06-06
; NUMBER OF SEQ ID NOS: 43
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 42
; LENGTH: 394
; TYPE: PRT
; ORGANISM: Triticum aestiva
US-09-424-978B-42

Query Match 90.8%; Score 1891.5; DB 4; Length 394;
Best Local Similarity 90.8%; Pred. No. 8.7e-192;
Matches 355; Conservative 18; Mismatches 17; Indels 1; Gaps 1;
QY 2 AETFLFTSESNEGHPDKLDCQISDAVLDAACLEQDPDSKVACETCTKTNLVMVFGIITK 61
Db |||||
QY 3 AETFLFTSESNEGHPDKLDCQISDAVLDAACLEQDPDSKVACETCTKTNLVMVFGIITK 62
Db |||||
QY 62 ANVDEYKIVRTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGHHTKRPBEIGAG 121
Db |||||
QY 63 ANVDEYKIVRTCRNIGFVSNVGLDADNCKVLNIEQQSPDIAQGVHGHHTKRPBEIGAG 122
Db |||||
QY 122 DQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGKTQVTVVEYNDN 181
Db |||||
QY 123 DQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNKGTCPWLRPDGKTQVTVVEYNDN 182
Db |||||
QY 182 GAMPVRVHTVLTSTQHDFTVNDIEAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 241
Db |||||
QY 183 GAMPVRVHTVLTSTQHDFTVNDIEAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFV 242
Db |||||
QY 242 GPHGDAGLTKRKIIIDTYGCGAHGGGAFSGKDPKVDKRGAYIVRQAAKSIIVASGLAR 301
Db |||||
QY 243 GPHGDAGLTKRKIIIDTYGCGAHGGGAFSGKDPKVDKRGAYIVRQAAKSIIVASGLAR 302
Db |||||
QY 302 RCIVQSVYAIGVPEPLSVFVDTYGTGKHDKHEILNIVKENFDFRPGMISINLDLKRGN 361
Db |||||
QY 303 RCIVQSVYAIGVPEPLSVFVDTYGTGKHDKHEILNIVKENFDFRPGMISINLDLKRGN 361
Db |||||
QY 362 RPLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db |||||
QY 362 RPLKTAAYGHGREDPDFTWEVVKPLKWEKA 392
Db |||||

RESULT 4
US-09-976-594-471
; Sequence 471, Application US/09976594
; Patent No. 6673549
; GENERAL INFORMATION:
; APPLICANT: Furness, Michael
; APPLICANT: Buchbinder, Jenny
; TITLE OF INVENTION: GENES EXPRESSED IN C3A LIVER CELL CULTURES TREATED WITH STEROIDS
; FILE REFERENCE: PA-0041 US
; CURRENT APPLICATION NUMBER: US/09/976,594
; CURRENT FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: 60/240,409
; PRIOR FILING DATE: 2000-10-12
; NUMBER OF SEQ ID NOS: 1143
; SOFTWARE: PERL Program
; SEQ ID NO 471
; LENGTH: 395
; TYPE: PRT
; ORGANISM: Homo sapiens

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;
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No. 6673549 2600262CD1
US-09-976-594-471

Query Match      63.2%; Score 1316.5; DB 4; Length 395;
Best Local Similarity 65.8%; Pred. No. 9.2e-131;
Matches 254; Conservative 49; Mismatches 76; Indels 7; Gaps 3;

QY 4 TELFTSESVEGHPDKLCQDISDAVLDALEQDPDSKVACETCTKTNLVVVGFEITTKAN 63
DB 17 TELFTSESVEGHPDKLCQDISDAVLDAHLQDPDAKACETVAKTGMILLAGEITSRAA 76

QY 64 VDYKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGV--HLDRNEEDIGAGD 123
DB 77 VDYQKVVREAVKHIGYDDSSKGFYDKTCNVLVVALEQQSPDIAQGV--HLDRNEEDIGAGD 134

QY 124 QGHMFGYATDETEPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYNDNGA 183
DB 135 QGLMFGYATDETEECMPLTIVLAHKNLAKLAELRRNGTLPWLRPDSKTQVTVQYMDRGA 194

QY 184 MYPVRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFVIGG 243
DB 195 VLPVRVHTVIVSQHDEEVCLEMDRALKEKVIKAVVPAKYLDDEDTIYHLQPSGRFVIGG 254

QY 244 PHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPYTKVDKRSAGYIVRQAASIVASGLARRC 303
DB 255 PQGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDYTKVDKRSAAAYAAARWAKSLVKGGLCRV 314

QY 304 IQVQSYAIGVPEPLSVFVDYTGTKIHDKEILNIVKENFDFRPGMISINLDLKRGGNNRF 363
DB 315 LVQVSYAIGVSHPLSISIFHYGTSQKSERELLEIVKQNFDLRPGVIVRDLDLKK---PIY 371

QY 364 LKTAAYGHGREDPDTWEVVKPLKW 389
DB 372 QRTAAYGHGFRD--SFPWEVPKPKLY 395

RESULT 5
US-09-949-016-10059
; Sequence 10059, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10059
; LENGTH: 416
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-10059

Query Match      63.2%; Score 1316.5; DB 4; Length 416;
Best Local Similarity 65.8%; Pred. No. 1e-130;
Matches 254; Conservative 49; Mismatches 76; Indels 7; Gaps 3;

QY 4 TELFTSESVEGHPDKLCQDISDAVLDALEQDPDSKVACETCTKTNLVVVGFEITTKAN 63
DB 38 TELFTSESVEGHPDKLCQDISDAVLDAHLQDPDAKACETVAKTGMILLAGEITSRAA 97

QY 64 VDYKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGV--HLDRNEEDIGAGD 123
DB 77 VDYQKVVREAVKHIGYDDSSKGFYDKTCNVLVVALEQQSPDIAQGV--HLDRNEEDIGAGD 134
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DB 98 VDYQKVVREAVKHIGYDDSSKGFYDKTCNVLVVALEQQSPDIAQGV--HLDRNEEDIGAGD 155
QY 124 QGHMFGYATDETEPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYNDNGA 183
DB 156 QGLMFGYATDETEECMPLTIVLAHKNLAKLAELRRNGTLPWLRPDSKTQVTVQYMDRGA 215
QY 184 MYPVRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFVIGG 243
DB 216 VLPVRVHTVIVSQHDEEVCLEMDRALKEKVIKAVVPAKYLDDEDTIYHLQPSGRFVIGG 275

QY 244 PHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPYTKVDKRSAGYIVRQAASIVASGLARRC 303
DB 276 PQGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDYTKVDKRSAAAYAAARWAKSLVKGGLCRV 335

QY 304 IQVQSYAIGVPEPLSVFVDYTGTKIHDKEILNIVKENFDFRPGMISINLDLKRGGNNRF 363
DB 336 LVQVSYAIGVSHPLSISIFHYGTSQKSERELLEIVKQNFDLRPGVIVRDLDLKK---PIY 392

QY 364 LKTAAYGHGREDPDTWEVVKPLKW 389
DB 393 QRTAAYGHGFRD--SFPWEVPKPKLY 416

RESULT 6
US-09-248-796A-18255
; Sequence 18255, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstein et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICA
; FILE REFERENCE: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 18255
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-18255

Query Match      61.5%; Score 1280.5; DB 4; Length 390;
Best Local Similarity 62.5%; Pred. No. 6e-127;
Matches 242; Conservative 56; Mismatches 82; Indels 7; Gaps 3;

QY 3 ETFLFTSESVEGHPDKLCQDISDAVLDALEQDPDSKVACETCTKTNLVVVGFEITTKA 62
DB 11 ETFLFTSESVEGHPDKLCQDISDAVLDALEQDPDSKVACETCTKTNLVVVGFEITTKA 70

QY 63 NVDEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVHGLTKRPEEIGAG 122
DB 71 QLDYQKIIIRDTIKHIGYDDSEKGFYDKTCNVLVVALEQQSPDIAQGL--HYEKALBELGAG 128

QY 123 QGHMFGYATDETEPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYNDNG 182
DB 129 QGHMFGYATDETEPELMPLSHVLATKLGARLTVVRKNGTCPLWRPDGKTQVTVVEYNDNG 188

QY 183 AMVPVRVHTVLISTQHDVTNDEIAADLKEHVIKPVIPEKYLDKTIIFHLNPSGRFVIG 242
DB 189 AVIPKRVDTIVISTQHAEEITENLRKEIIIEHIIKQVPEHLDDKTIYHIQPSGRFVIG 248

QY 243 GPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPYTKVDKRSAGYIVRQAASIVASGLARR 302
DB 249 GPQGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPYTKVDKRSAAAYAAARWAKSLVTAGLAKR 308

QY 303 CIVQVSYAIGVPEPLSVFVDYTGTKIHDKEILNIVKENFDFRPGMISINLDLKRGGNNR 362
DB 309 ALVQPSYAIGVAEPTSIYIDTYGTSKLSLEALVEIIKKNFDLRPGVIVKELDLAR---PI 365
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; APPLICATION NUMBER: 60/085,598
; FILING DATE: 14 May 1998
; APPLICATION NUMBER: 60/051571
; FILING DATE: July 2, 1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Ariniello, Pamela Deneke
; REGISTRATION NUMBER: 40,489
; REFERENCE/DOCKET NUMBER: GTC-012
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (781)893-5007
; TELEFAX: (781)893-8277
; INFORMATION FOR SEQ ID NO: 6821:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 404 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; HYPOTHEICAL: YES
; ORIGINAL SOURCE:
; ORGANISM: Enterococcus faecium
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (B) LOCATION 1...404
; SEQUENCE DESCRIPTION: SEQ ID NO: 6821:
US-09-107-532A-6821

Query Match      52.9%; Score 1102; DB 4; Length 404;
Best Local Similarity 57.2%; Pred. No. 5.6e-108;
Matches 223; Conservative 60; Mismatches 93; Indels 14; Gaps 5;

QY 1 MAETFLFTSSEVNEGHPD-KLCDQISDAVLDACLEQDPDSKVACETCTKTNLVNVFGEIT 59
DB 7 MVERHLFTSSEVSEHPRIKADQISDAVLDAILDKQDPTARVACETSVTTGLVLFGEIS 66

QY 60 TKANVDYEKIVRTCRNIGFVSNVDGLDADNCKVLNIEQSPDIAQGVHGLTKRPE-- 117
DB 67 TTAVIDIQVRETIKEIGYTRAKFGDGTAAVLVAIDEQSPDIAQGVDEALEIRDEDK 126

QY 118 -----BIGAGDQGHMFGYATDETPELMPLSHVLATKLGARLTVEVRKNGTCPLRPDGKTQ 172
DB 127 KOVLDEIGAGDQGLMFGFAVDETPELMPLSHVLRLADLRKSNELTYLRPDASKQ 186

QY 173 VTVEYNDNGAMVPVRVHTVLTSTQHDENVNDEIAADLKEHVIKPVIPEKYLDEKTI 232
DB 187 VTVE-YDDQ--PERVDTIVISTQHDADVNETIRHVDIEKVKEVIPAELLDDQTKY 243

QY 233 LNPGRFVIGGPHGDAGLTKRKIIIDTYGNGWAGHGGAFSGKDPKTVDRSGAYIVRQAAK 292
DB 244 INPTGRFVIGGPGQDAGLTKRKIIIDTYGNGWAGHGGAFSGKDATKVDRSASAYARIYAK 303

QY 293 SIVASGLARRCIVQVSYAIGVPEPLSVFVDYTGTKIHDKELINIVKENFDRPFGMISIN 352
DB 304 NIVAAGLARKAEVQLAYAGVQVPSISINTFGTVPEELAAVRENFDLRPAGIIE 363

QY 353 LDLKRGNNRFLKTAAYGHGREDPDFTWE 382
DB 364 LDLRR---PIYKTAAYGHGFRDVLPMWE 390

RESULT 10
US-09-543-681A-7130
; Sequence 7130, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709, 1002-001
; CURRENT APPLICATION NUMBER: US/09/543, 681A
; PRIOR FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
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; SEQ ID NO 7130
; LENGTH: 387
; TYPE: PRT
; ORGANISM: Proteus mirabilis
; US-09-543-681A-7130

Query Match      52.1%; Score 1086; DB 4; Length 387;
Best Local Similarity 58.2%; Pred. No. 2.6e-106;
Matches 221; Conservative 56; Mismatches 89; Indels 14; Gaps 6;

QY 4 TFLFTSSEVNEGHPDKLCDQISDAVLDACLEQDPDSKVACETCTKTNLVNVFGEITTKAN 63
DB 6 TFLFTSSEVSEGHDPKIDQISDAVLDAILDQDPKARVACETVYVTKGMVVGGEITTKAW 65

QY 64 VDEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSPDIAQGVHGLTKRPEICAGD 123
DB 66 VDIEITRTVREIGYSSDMGFDANCAVISAIGKQSPDINQGV--RADPLEQAGD 122

QY 124 QGHMFGYATDETPELMPLSHVLATKLGARLTVEVRKNGTCPLRPDGKTQTVVEYNDNGA 183
DB 123 QGLMFGYATNETDVLMPAPITVAHRLVQROAQVRKNGTLPWLRPDASKQITTFQYDNN-- 180

QY 184 MYPVRVHTVLTSTQHDENVNDEIAADLKEHVIKPVIPEKYLDEKTIPLHNSGRFVIG 243
DB 181 --IVGIDAVLSTQHAEDISQDLHEAVMBEIIKPLPTEWLNQTKYFINPTGRFVIG 238

QY 244 PHGDAGLTKRKIIIDTYGNGWAGHGGAFSGKDPKTVDRSGAYIVRQAAKSIVASGLARRC 303
DB 239 PMGDCGLTKRKIIIDTYGNGWAGHGGAFSGKDPKTVDRSAAVYAKNIIVAAGLADRC 298

QY 304 IVQVSYAIGVPEPLSVFVDYTGTKIHDKELINIVKENFDRPFGMISINLDKRGNNR 362
DB 299 EIQVSYAIGVPEPLSVFVDYTGTKIHDKELINIVKENFDRPFGMISINLDKRGNNR 362

QY 363 FLKTAAYGHGREDPDFTWE 382
DB 355 YOKTAAYGHGFR--AEFPWE 372

RESULT 11
US-09-489-039A-11917
; Sequence 11917, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et. al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; FILE REFERENCE: 2709, 2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 11917
; LENGTH: 385
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
; US-09-489-039A-11917

Query Match      51.3%; Score 1068.5; DB 4; Length 385;
Best Local Similarity 57.5%; Pred. No. 1.9e-104;
Matches 223; Conservative 54; Mismatches 92; Indels 19; Gaps 8;

QY 6 LFTSSEVNEGHPDKLCDQISDAVLDACLEQDPDSKVACETCTKTNLVNVFGEITTKANVD 65
DB 6 LFTSSEVSEGHDPKIDQISDAVLDAILDQDPKARVACETVYVTKGMVVGGEITTSWVD 65

QY 66 YEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSPDIAQGVHGLTKRPEICAGDQ 125
DB 66 IBEITRTVREIGYSSDMGFDANCAVISAIGKQSPDINQGV--RADPLEQAGDQ 122

QY 126 HMFYATDETPELMPLSHVLATKLGARLTVEVRKNGTCPLRPDGKTQTVVEYNDNGAMV 185
```

Db 123 LMFYATNETDVLMPAPVTYAHRLVQROAEVRKNGTLPWLDPDAKSQVTFQY--DDGKI- 179

QY 186 PVRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTTIPLNPSGRFVIGGPH 245

Db 180 -VGIDAVVLSTQHAEDIDQKSLQEAWMEEIIPILPTWLNASTKFFINPTGRFVIGGPM 238

QY 246 DGAGLTGRKIIIDTGGWGAHGGGAFSGDKPTKVDRSYIVROAKSVASGLARRCIV 305

Db 239 GDCGLTGRKIIIDTGGWGAHGGGAFSGDKPSKVDRSAAAYAAKYAKNIVAAGLADRCI 298

QY 306 QVSYAIGVPEPLSVFVDYTGTKIHDKEILNIVKENFDRP-GMISINLDLXRGGNRFL 364

Db 299 QVSYAIGVAEPTSIMVETGTETKVPSEQLTLVREFFDLRPYGLIQM-LDLL--HPIYK 354

QY 365 KTAAYGHGREDDPDTFWEVVKPLKWEKA 392

Db 355 ETAAYGHGRE--HFPWE-----KTDKA 375

RESULT 12

US-09-583-110-2778

; Sequence 2778, Application US/09583110

; Patent No. 6699703

; GENERAL INFORMATION:

; APPLICANT: Lynn Doucette-Stamm et al.

; TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus

; FILE REFERENCE: PATH00-07A

; CURRENT APPLICATION NUMBER: US/09/583,110

; PRIOR FILING DATE: 2000-05-26

; PRIOR APPLICATION NUMBER: US 09/107,433

; PRIOR FILING DATE: 1998-06-30

; PRIOR APPLICATION NUMBER: US 60/085,131

; PRIOR FILING DATE: 1998-05-12

; PRIOR APPLICATION NUMBER: US 60/051,553

; PRIOR FILING DATE: 1997-07-02

; NUMBER OF SEQ ID NOS: 5322

; SEQ ID NO 2778

; LENGTH: 396

; TYPE: PRT

; ORGANISM: Streptococcus pneumoniae

US-09-583-110-2778

Query Match 51.0%; Score 1062.5; DB 4; Length 396;

Best Local Similarity 55.9%; Pred. No. 8.4e-104;

Matches 218; Conservative 58; Mismatches 99; Indels 15; Gaps 5;

QY 1 MAETFLFTSESVEGHGPKLDCQISDAVLDALEQDPDSKVACETCTKTNLVMVFGEIT 60

Db 1 MSERKLFTSSEVSEGHGPKIADQISDAILDALAKDPEAHVAEAETAVYTGSVHVFGEIST 60

QY 61 KANVDYKIVRDTCTRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVH-----GHLTK 114

Db 61 NAVVDINRVVRDITAEIGYTNTEYGFSAETGVVHPSLVESQSPDIAQGVNEALEVRGNADQ 120

QY 115 RP-EBIGAGDQGHMFYATDETPELMPLSHVLATKLGARLTVRNKGTCPWLRPDGKTQV 173

Db 121 DPLDLIAGDQGLMFGFAVDETEELMPLIALSHKLVRLAELRKSGEISYLRPDAKSQV 180

QY 174 TVEY-YNDNGAMPVVRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTTI 232

Db 181 TVEYDENDR-----PVRVTVTVLSTQHDPEATNEQIHQDVIDKVIKEVIPSLLDDKTKFF 236

QY 233 LNPGRFVIGGPHDAGLTGRKIIIDTGGWGAHGGGAFSGDKPTKVDRSYIYVQAAK 292

Db 237 INPTGRFVIGGPQDGLTGRKIIIDTGGYSRHGGGAFSGDKATKVDRSASVAAARYIAK 296

QY 293 SIVASGLARRCIVQVSYAIGVPEPLSVFVDYTGTKIHDKEILNIVKENFDRPGRMISIN 352

Db 297 NIYAAGLAKAEVQLAYAIGVAQVPSVRIDTFGTGTVAESQLEKAAARQIFDLRPAGIIQM 356

QY 353 LDLKRGNNRFLKTAAYGHGREDDPDTWE 382

Db 357 LDLKR---PIYRQTSAYGHMGRGTIDILPWE 383

RESULT 13

US-09-107-433-2830

; Sequence 2830, Application US/09107433

; Patent No. 6800744

; GENERAL INFORMATION:

; APPLICANT: Lynn A Doucette-Stamm and David Bush

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE THERAPEUTICS

; NUMBER OF SEQUENCES: 5206

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: GENOME THERAPEUTICS CORPORATION

; STREET: 100 Beaver Street

; CITY: Waltham

; STATE: Massachusetts

; COUNTRY: USA

; ZIP: 02354

; COMPUTER READABLE FORM:

; MEDIUM TYPE: CD-ROM ISO9660

; COMPUTER: <Unknown>

; OPERATING SYSTEM: <Unknown>

; SOFTWARE: <Unknown>

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/09/107,433

; FILING DATE: 30-Jun-1998

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 60/ 085131

; FILING DATE: May 12, 1998

; APPLICATION NUMBER: 60/051553

; FILING DATE: July 2, 1997

; ATTORNEY/AGENT INFORMATION:

; NAME: Ariniello, Pamela Deneke

; REGISTRATION NUMBER: 40,489

; REFERENCE/DOCKET NUMBER: GTC-011

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: (781)893-5007

; TELEFAX: (781)893-8277

; INFORMATION FOR SEQ ID NO: 2830:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 405 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; HYPOTHETICAL: YES

; ORIGINAL SOURCE:

; ORGANISM: Streptococcus pneumoniae

; FEATURE:

; NAME/KEY: misc feature

; LOCATION: (B) LOCATION 1...405

; SEQUENCE DESCRIPTION: SEQ ID NO: 2830:

US-09-107-433-2830

Query Match 51.0%; Score 1062.5; DB 4; Length 405;

Best Local Similarity 55.9%; Pred. No. 8.7e-104;

Matches 218; Conservative 58; Mismatches 99; Indels 15; Gaps 5;

QY 1 MAETFLFTSESVEGHGPKLDCQISDAVLDALEQDPDSKVACETCTKTNLVMVFGEIT 60

Db 10 MSERKLFTSSEVSEGHGPKIADQISDAILDALAKDPEAHVAEAETAVYTGSVHVFGEIST 69

QY 61 KANVDYKIVRDTCTRNIGFVSNVDGLDADNCKVLNIEQQSPDIAQGVH-----GHLTK 114

Db 70 NAVVDINRVVRDITAEIGYTNTEYGFSAETGVVHPSLVESQSPDIAQGVNEALEVRGNADQ 129

QY 115 RP-EBIGAGDQGHMFYATDETPELMPLSHVLATKLGARLTVRNKGTCPWLRPDGKTQV 173

Db 130 DPLDLIAGDQGLMFGFAVDETEELMPLIALSHKLVRLAELRKSGEISYLRPDAKSQV 189

QY 174 TVEY-YNDNGAMPVVRVHTVLSTQHDFTVTNDEIAADLKEHVIKPVIPEKYLDKTTI 232

Db 190 TVEYDENDR----PVRVDTVVIQTQHDPEATNEQIHQVDIKVIKEVIPSSYLDDTKTFF 245
QY 233 LNPSGRFVIGGPHDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKTVDRSGAYIVRQAQ 292
Db 246 INPTGRFVIGGPGDGLTKRKIIIDTYGGYSHGGGAFSGKDPKTVDRSGAYIVRQAQ 305
QY 293 SIVASGLARRCIVQSVYAIGVPEPLSVFVDTYGTGKHDKKEILNIVKENFDFRPGMISIN 352
Db 306 NIVAAGLAKAEVQLAYAIGVAPVSVRIIDTGTGTVAESQLEKAAQIFDLRPAGLIQM 365
QY 353 LDLKRGNNRFLKTAAYGHGREDPDFTWE 382
Db 366 LDLKRL----PIYRQTSAYGHMGRTDIDLPE 392

RESULT 14

US-09-273-686-2

; Sequence 2, Application US/09273686

; Patent No. 6228625

; GENERAL INFORMATION:

; APPLICANT: Zalacain, Magdalena

; APPLICANT: Burnham, Martin K. R.

; APPLICANT: Biswas, Sanjoy

; APPLICANT: Brown, James

; APPLICANT: Ingraham, Karen, A.

; APPLICANT: Chalker, Alison F.

; APPLICANT: So, Chi Y.

; APPLICANT: Holmes, David J.

; APPLICANT: Van Horn, Stephanie

; APPLICANT: Warren, Richard L.

; FILE REFERENCE: metK

; CURRENT APPLICATION NUMBER: US/09/273,686

; CURRENT FILING DATE: 1999-03-22

; EARLIER APPLICATION NUMBER: 60/106,767

; EARLIER FILING DATE: 1998-11-03

; NUMBER OF SEQ ID NOS: 3

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 2

; LENGTH: 396

; TYPE: PRF

; ORGANISM: Streptococcus pneumoniae

US-09-273-686-2

Query Match 51.0%; Score 1061.5; DB 3; Length 396;

Best Local Similarity 55.9%; Pred. No. 1.1e-103;

Matches 218; Conservative 58; Mismatches 99; Indels 15; Gaps 5;

QY 1 MAETFLTSESNEGHDPKLCDOISDAVLDALEQDPPDSKVACETCTKTNLMVVFGEIT 60
Db 1 MSERKLTSESNEGHDPKLCDOISDAVLDALEQDPPDSKVACETCTKTNLMVVFGEIT 60
QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSPDIAQGVH-----GHLTK 114
Db 61 NAVVDINRVVDTIAEIGYNTTEYGFSAETGVHPSLVEQSPDIAQGVNEALEVRGNADQ 120
QY 115 RP-EEIGAGQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNGTCPWLRPDQKTV 173
Db 121 DPLDLIGAGDQGLMFGFVADTEBELMPLSHVLATKLGARLTVRKNGTCPWLRPDQKTV 180
QY 174 TVEY-YNDNGAMPVVRVHTVLSTQHDPTVNDIAADLKEHVKIPVPEKYLDEKTIHF 232
Db 181 TVEYDENDR----PVRVDTVVIQTQHDPEATNEQIHQVDIKVIKEVIPSSYLDDTKTFF 236
QY 233 LNPSGRFVIGGPHDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKTVDRSGAYIVRQAQ 292
Db 237 INPTGRFVIGGPGDGLTKRKIIIDTYGGYSHGGGAFSGKDPKTVDRSGAYIVRQAQ 296
QY 293 SIVASGLARRCIVQSVYAIGVPEPLSVFVDTYGTGKHDKKEILNIVKENFDFRPGMISIN 352
Db 297 NIVAAGLAKAEVQLAYAIGVAPVSVRIIDTGTGTVAESQLEKAAQIFDLRPAGLIQM 356
QY 353 LDLKRGNNRFLKTAAYGHGREDPDFTWE 382

Db 357 LDLKRL----PIYRQTSAYGHMGRTDIDLPE 383

RESULT 15

US-09-328-352-6660

; Sequence 6660, Application US/09328352

; Patent No. 6562958

; GENERAL INFORMATION:

; APPLICANT: Gary L. Breton et al.

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER

; FILE REFERENCE: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS

; CURRENT APPLICATION NUMBER: US/09/328,352

; CURRENT FILING DATE: 1999-06-04

; NUMBER OF SEQ ID NOS: 8252

; SEQ ID NO 6660

; LENGTH: 395

; TYPE: PRF

; ORGANISM: Acinetobacter baumannii

US-09-328-352-6660

Query Match 50.6%; Score 1055; DB 4; Length 395;

Best Local Similarity 54.9%; Pred. No. 5.2e-103;

Matches 211; Conservative 61; Mismatches 100; Indels 12; Gaps 5;

QY 1 MAETFLTSESNEGHDPKLCDOISDAVLDALEQDPPDSKVACETCTKTNLMVVFGEIT 60
Db 8 MREYAVFTSESNEGHDPKLCDOISDAVLDALEQDPPDSKVACETCTKTNLMVVFGEIT 67
QY 61 KANVDYEKIVRDTCRNIGFVSNVDGLDADNCKVLNIEQSPDIAQGVHGLTKRPEEIG 120
Db 68 TANIDVEAVRQTVNGIGYHSDLGFDGSCAVINMIGKQSPDIAQGVHGLTKRPEEIG 124
QY 121 AGDQGHMFGYATDETPELMPLSHVLATKLGARLTVRKNGTCPWLRPDQKTVTVYND 180
Db 125 AGDQGLMFGYASRETDLVMPAPISYAHRLMERQAEILRRSGALPWLRPDAKSQVTFAY--E 182
QY 181 NGAMPVVRVHTVLSTQHDPTVNDIAADLKEHVKIPVPEKYLDEKTIHFILNPSGRFV 240
Db 183 NGK--PVRLDVAVLSTQHDPTVNDIAADLKEHVKIPVPEKYLDEKTIHFILNPSGRFV 240
QY 241 IGGPHGDAGLTGRKIIIDTYGGWGAHGGGAFSGKDPKTVDRSGAYIVRQAQSIASGLA 300
Db 241 IGGPVGDCGLTKRKIIIDTYGGWGAHGGGAFSGKDPKTVDRSGAYIVRQAQSIASGLA 300
QY 301 RRCIVQSVYAIGVPEPLSVFVDTYGTGKHDKKEILNIVKENFDFRPGMISINLDKRGNN 360
Db 301 DKCEIQVSYAIGVAPVSVRIIDTGTGTVAESQLEKAAQIFDLRPAGLIQM--- 357
QY 361 NRFLKTAAYGHGREDPD--FTWE 382
Db 358 PMYKQTAAYGHGREGSDTAFTWE 381

Search completed: September 12, 2005, 20:56:54

Job time : 34 secs

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